IN THE SPECIFICATION

(1) At column 1, after the title, please insert the following sentence:

Notice: More than one reissue application has been filed for the reissue of Patent No. 5,691,454. The reissue applications are application numbers 09/442,489 (the present application) and 09/983,543, which is a continuation reissue of Patent No. 5,691,454.

(2) Delete the paragraph at column 4, lines 9-18, and replace it with the following paragraph:

FIGS. 2A and 2B show the sequence of TB1 (FIG. 2A, SEQ ID NO:5) and TB2 (FIG. 2B, SEQ ID NO:6) [genes] proteins. The cDNA sequence of the TB1 gene was determined from the analysis of 11 cDNA clones derived from normal colon and liver, as described in the text. A total of 2314 bp were contained within the overlapping cDNA clones, defining an ORF of 424 amino acids beginning at nucleotide 1. Only the predicted amino acids from the ORF are shown. The carboxy-terminal end of the ORF has apparently been identified, but the 5' end of the TB1 transcript has not yet been precisely determined.

(3) At column 32, delete Table IV and replace it with:

TABLE IV

Seven Different Versions of the 20-Amino Acid Repeat

(SEQ ID NO:147)	(SEQ ID NO:148)	(SEQ ID NO:149)	(SEQ ID NO:150)	(SEQ ID NO:151)	(SEQ ID NO:152)	(SEQ ID NO:153)	(SEQ ID NO:154)
F * V E * T P * C F S R * S S L S S L S	YCVEDTPICFSRCSSLSSLS	HTVQETPLMFSRCTSVSSLD	FATESTPDGFSCSSSLSALS	YCVEGTPINFSTATSLSDLT	TPIEGTPYCFSRNDSLSSLD	FAIENTPVCPSHNSSLSSLS	RHVEDTPVCFSRNSSLSSLS
Consensus:	1262:	1376:	1492:	1643:	1848:	1953:	2013:

Numbers denote the first amino acid of each repeat. The consensus sequence at the top reflects a majority amino acid at a given position. In the consensus sequence, "*" indicates "Xaa."

(4) At columns 31-132, please delete the sequence listing and substitute the following amended substitute sequence listing.